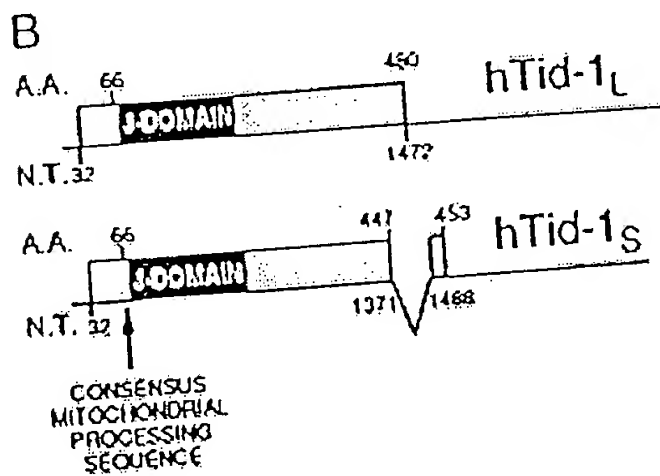
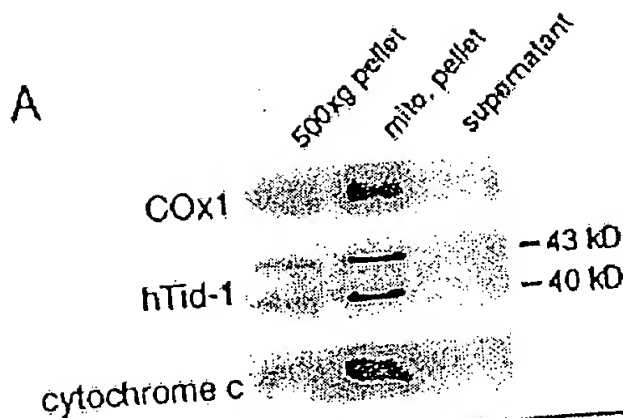
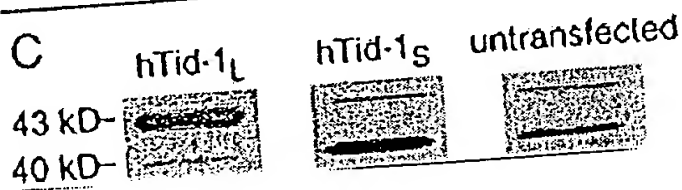


Figure 1



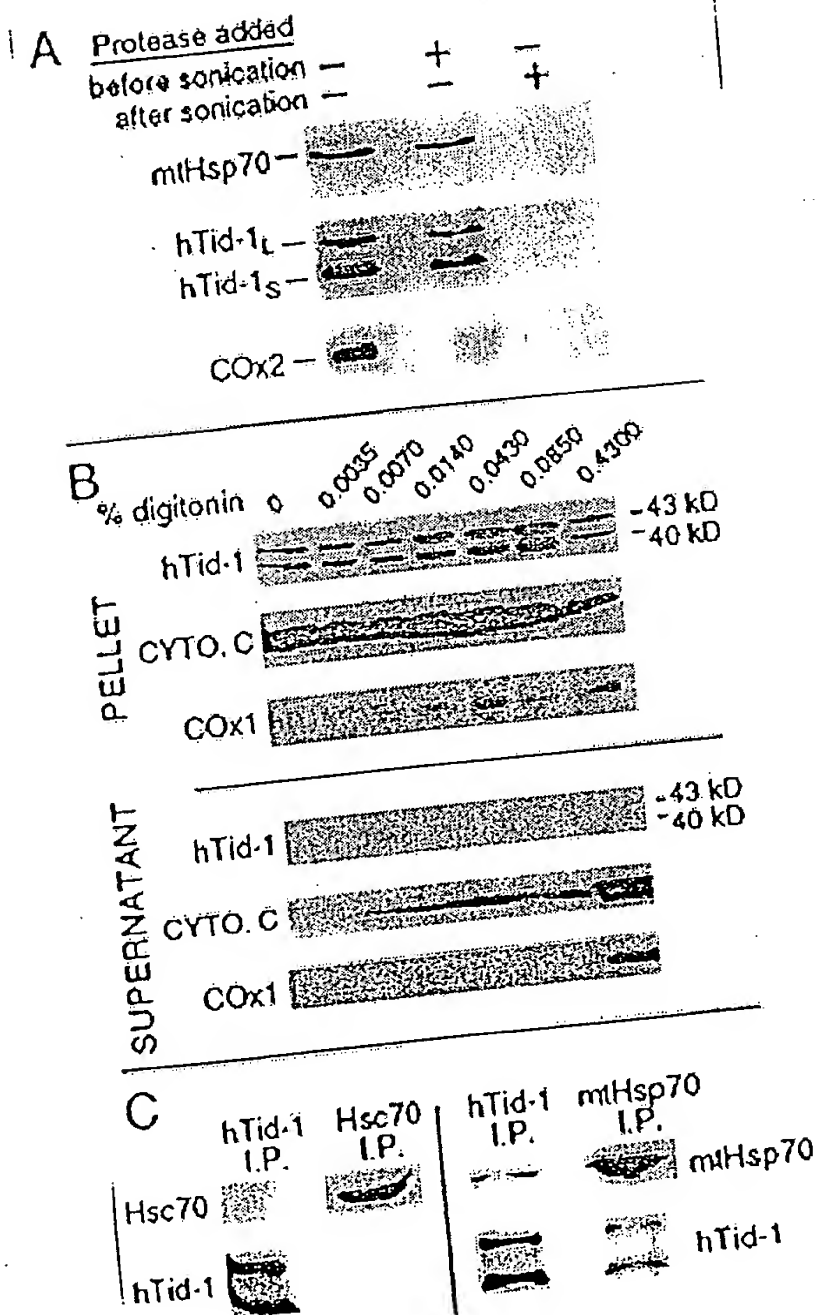
hTid-1_L carboxyl terminus
SSGGSTMDSAGSKARREAGEDEEGFLSKLKKMFTS

hTid-1_S carboxyl terminus
SSGKRSTGN



106720-26680660

Figure 2



106140-2660660

Figure 3

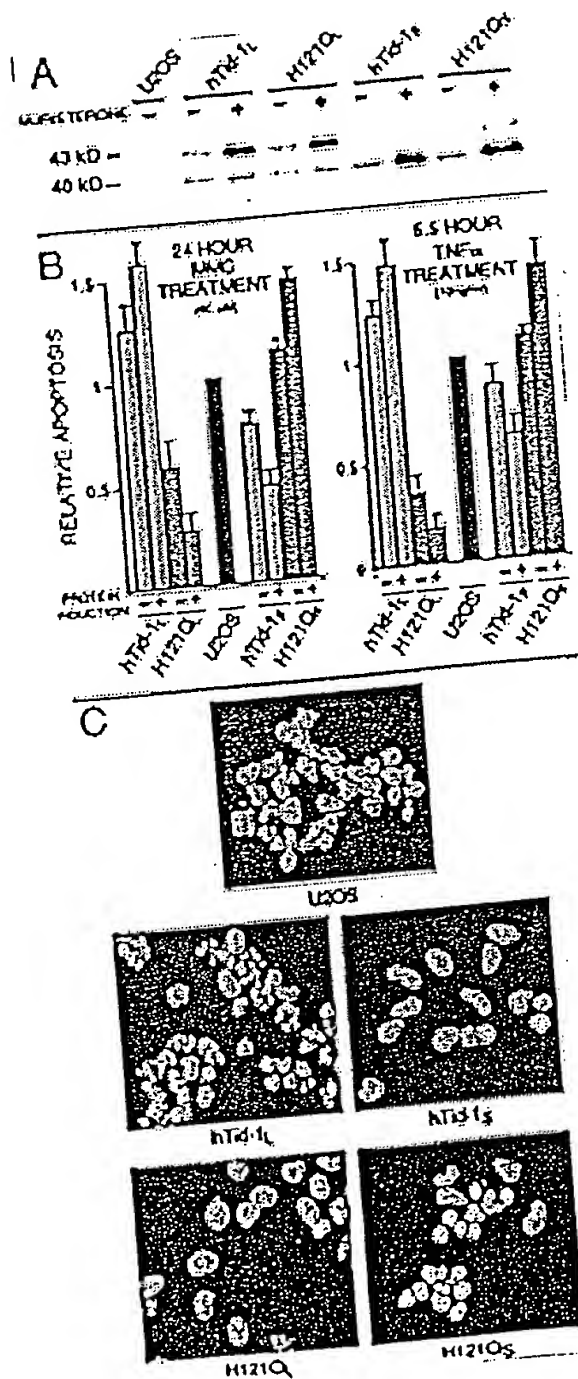
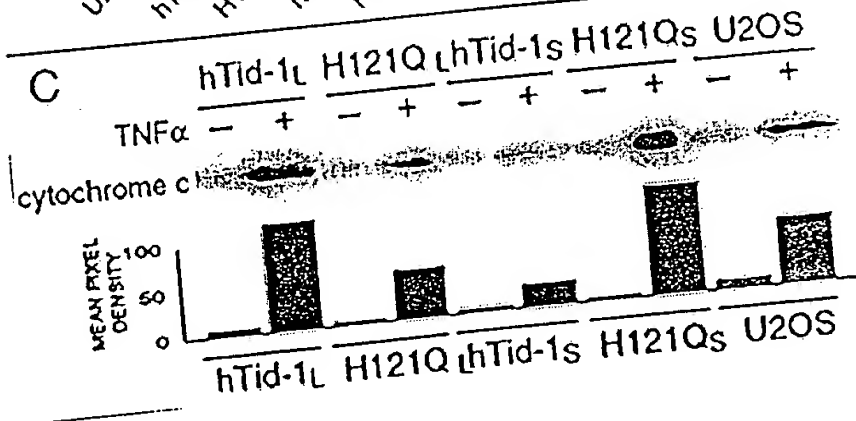
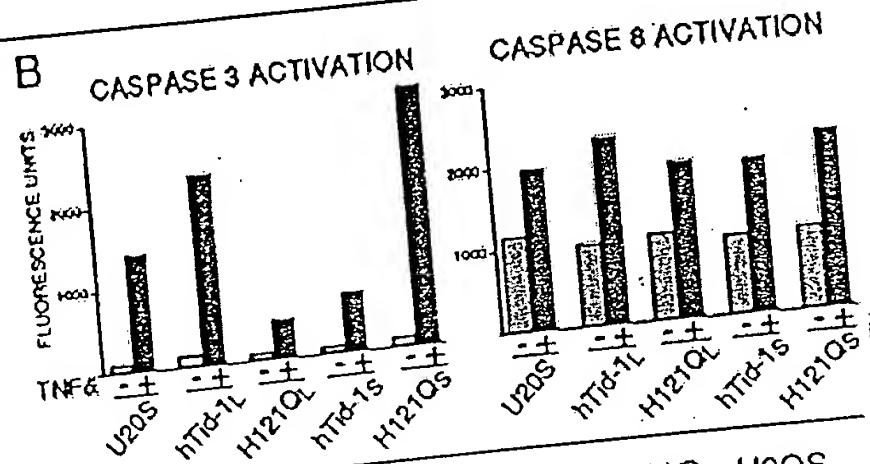
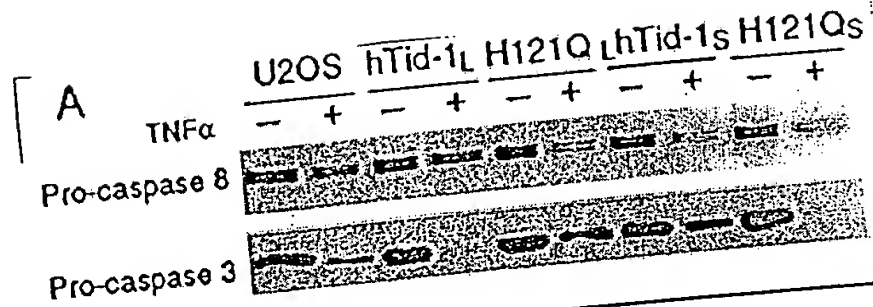


Figure 4



106720-20000000

Figure 5

Nucleotide sequence of coding sequence encoding hTid1L:

ATGgctgcgcggtgctccacacgctggttgcgtggtggttgcgtgggaccccgcggtgcccggctatatcgggtagaggggc
 ccggccgcccaggagggcggtggtgggggcatggctgagccgcaagctgagcgtccccgcctttgcgtcttccctgacct
 cttgcggccccgagcgctgctgacattgagacctggtgtcagccttacaggaacaaaacataaccctttcatttgtact
 gcctccttccacacgagtgcccccttggccaaagaagattattatcagatattaggagtgctcgaaatgccagccagaa
 agagatcaagaaagcctattatcagcttgccaagaagtatcacctgacacaaataaggatgatcccaaagccaaggaga
 agttctcccagctggcagaagcctatgaggttttgagtgatgaggtgaagaggaagcagtagcatgctacggctctgca
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 gaagatccttggcgagttctcatcctcttcatgttgagatttccagaccgtgttgatcagcctcaggaatacttcatgg
 agttgacattcaatcaagctgcaaaggggtcaacaaggagttcacctgaacatcatggacacgtgtgagcgctgcaac
 ggcaaggggaacgagcccgccaccaaggtgcagcattgccactactgtggcggtcccgcatggaaacctcaacacagg
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 tacggagaccactacatccacatcaagatacaggttccaaagaggctaaccgagcctcgggtggcagcaccatggatagct
 cgccgaggacgagacagatgtggaggggacgggtgaacggcgctcacccctcaccagctctgggtggcagcaccatggatagct
 ccgaggaagcaaggctaggcgtaggctggggaggacgaggaggatcccttccaaacttaagaaaatgtttacctca
 TGA

Nucleotide sequence of coding sequence encoding hTid1S:

atggctgcGcggtgctccacacgctggttgcgtggtggttgcgtgggaccccgcggtgcccggctatatcgggtagaggggc
 ccggccgcccaggagggcggtggtgggggcatggctgagccgcaagctgagcgtccccgcctttgcgtcttccctgacct
 cttgcggccccgagcgctgctgacattgagacctggtgtcagccttacaggaacaaaacataaccctttcatttgtact
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 agttctcccagctggcagaagcctatgaggttttgagtgatgaggtgaagaggaagcagtagcatgctacggctctgca
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 AG

FOR 266800

TOPFAD 26630660

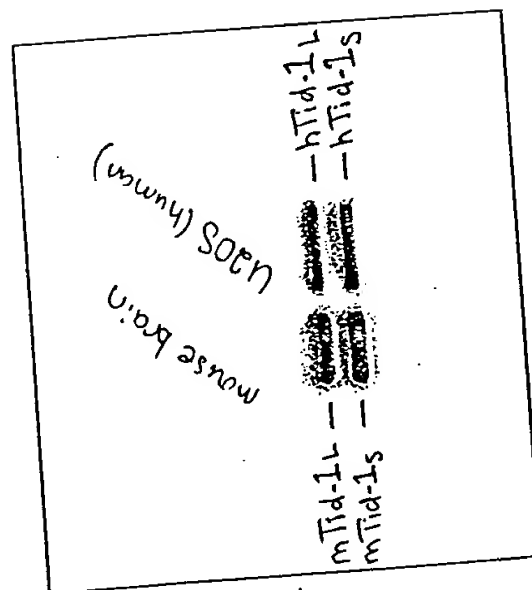
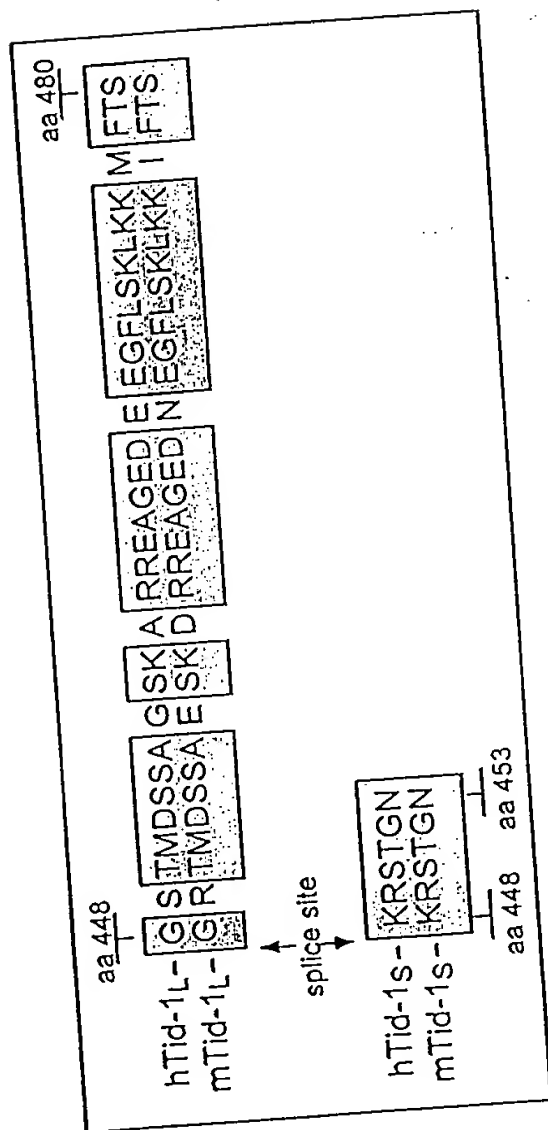


Figure 6

TOP 20 2668060

hrs activation

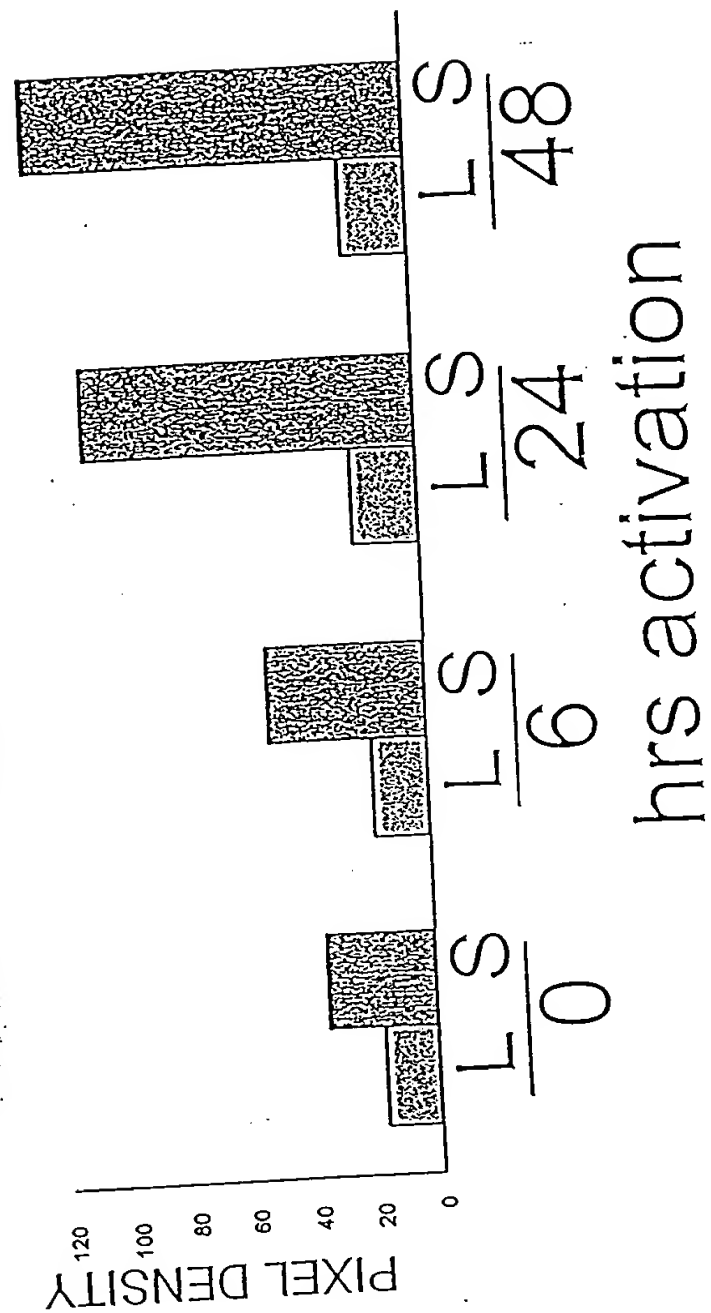
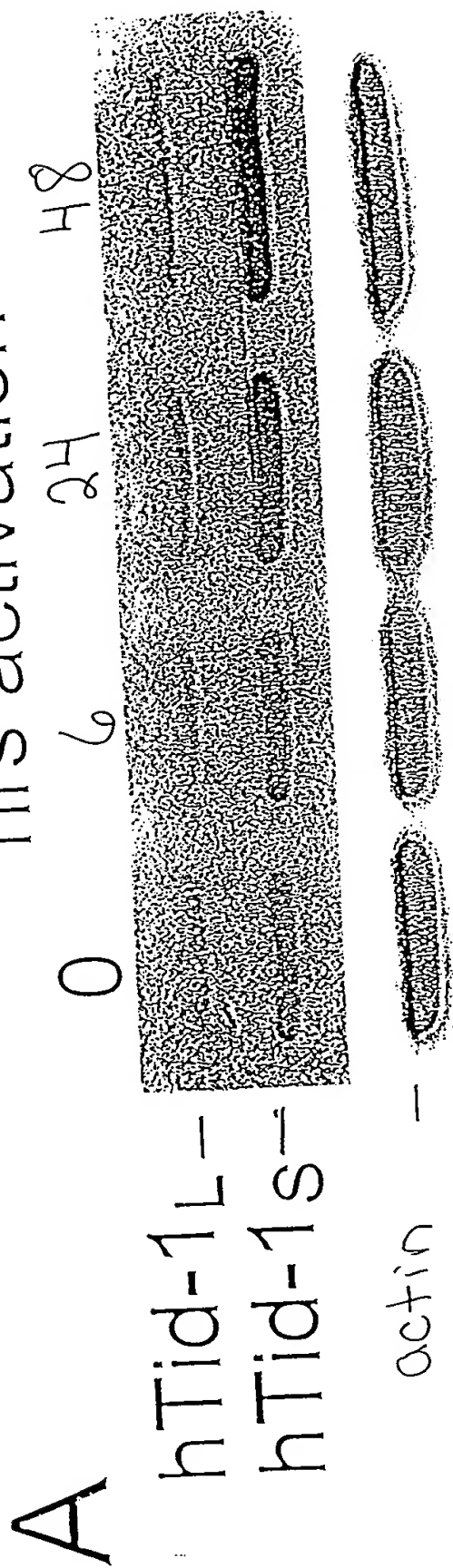


Figure 7A

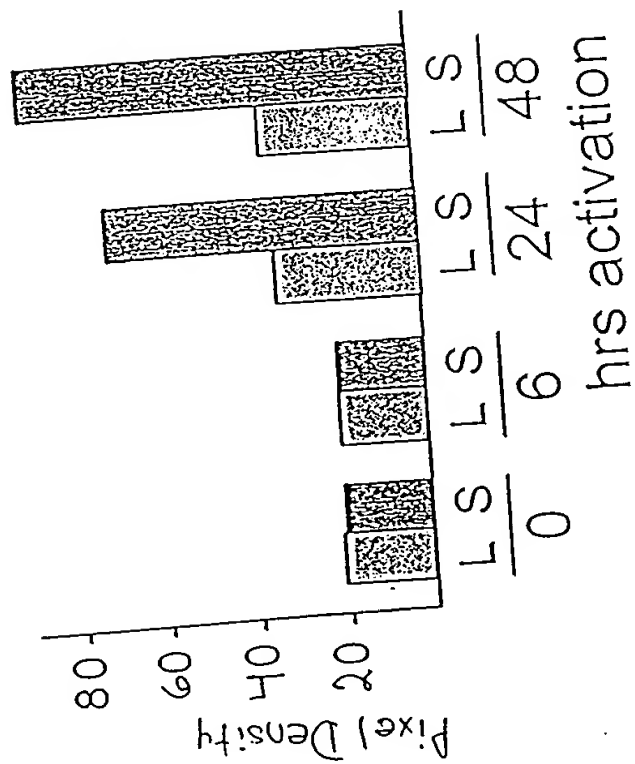
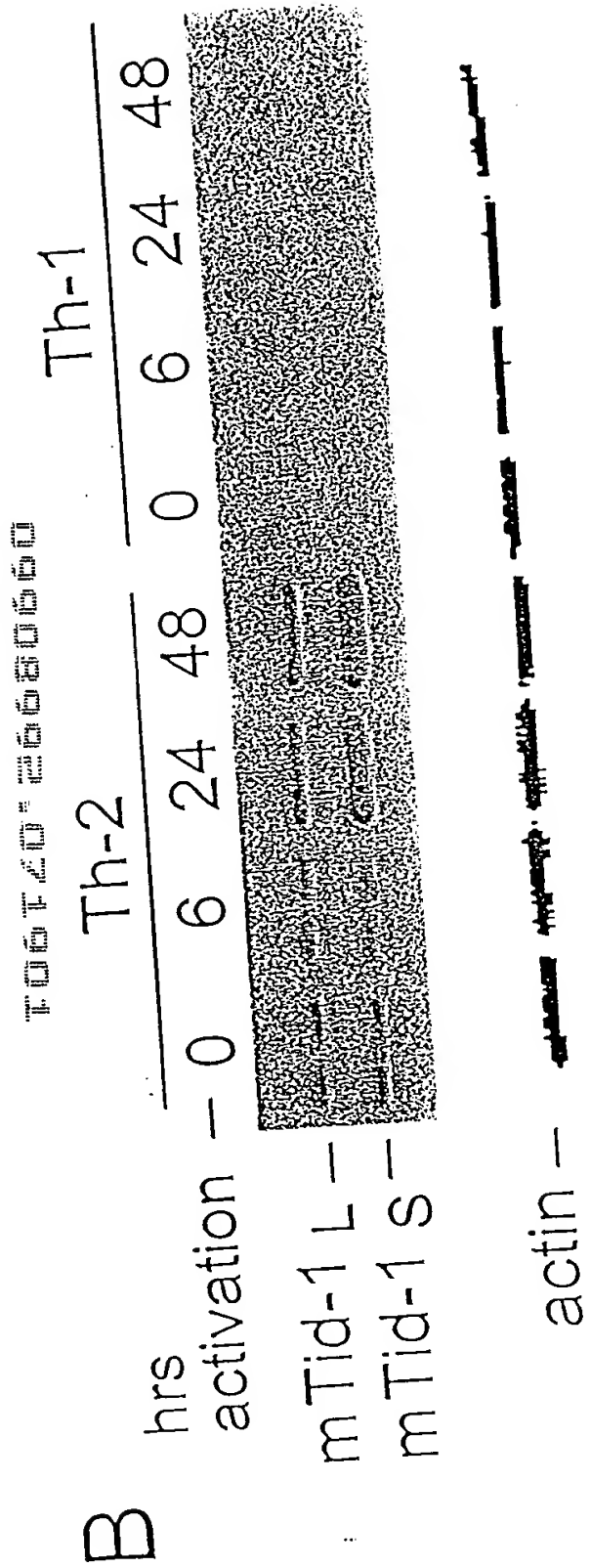


Figure 7B

T06T20" 2668060

C

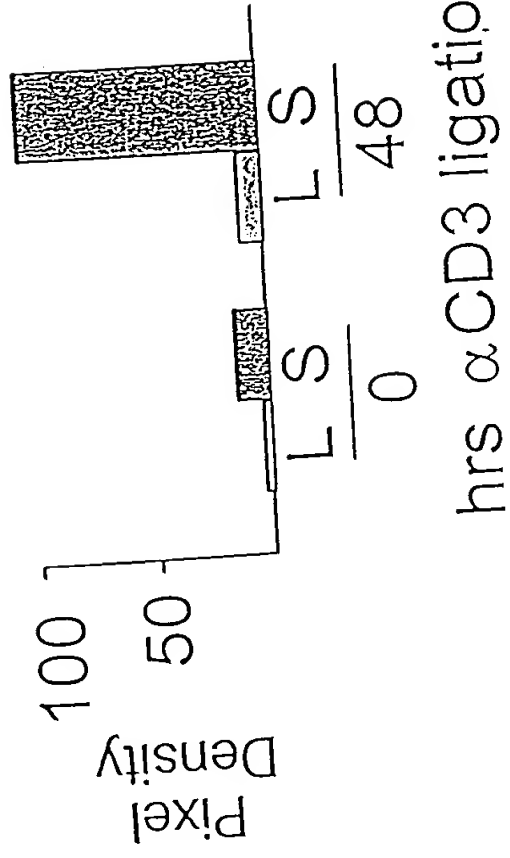
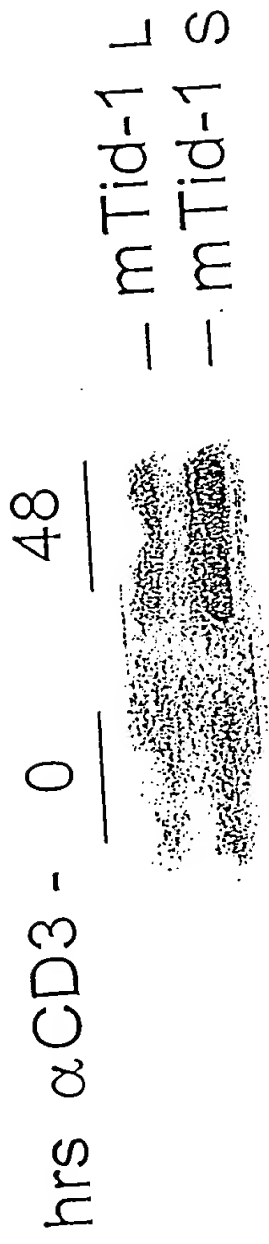


Figure 7C

106120 25680000

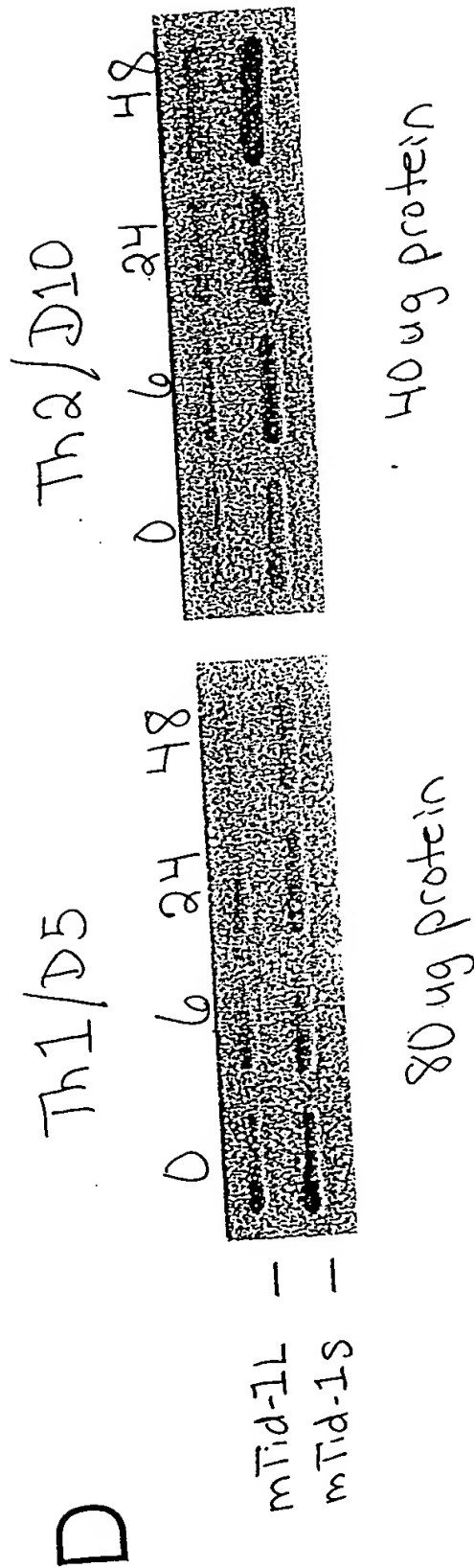


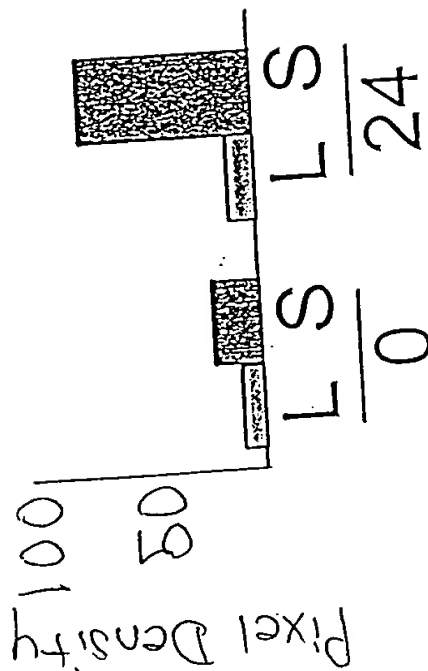
Figure 7D

W

 α_{CD3}

mTid-1L-

mTid-1s-



DO. 10. 11
primary Th2 cells

Figure 7E

TOPFLO 26680660

F

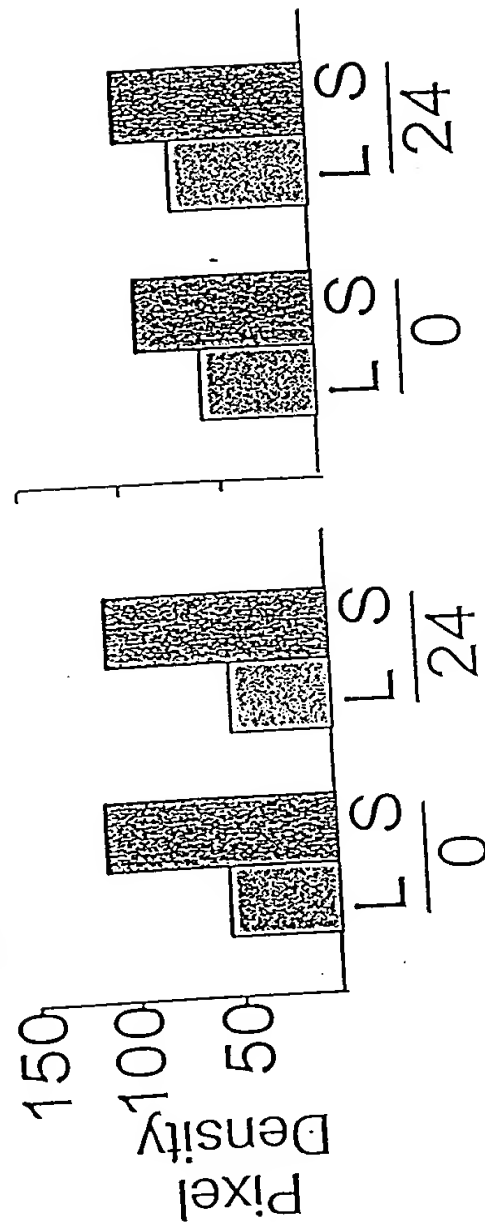
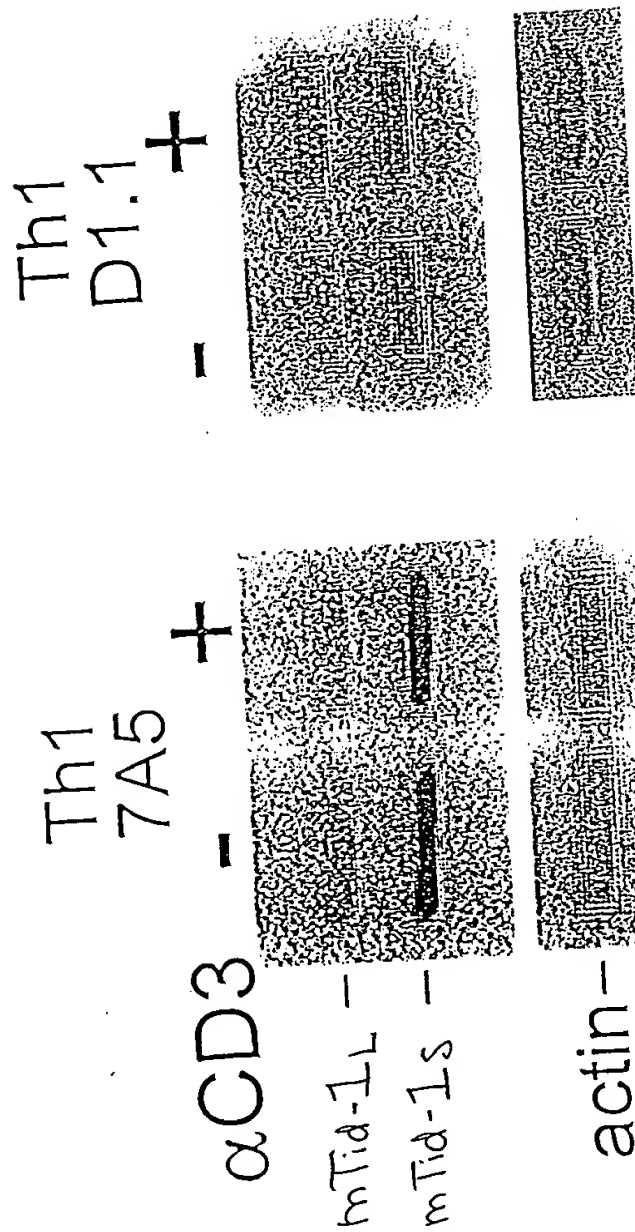


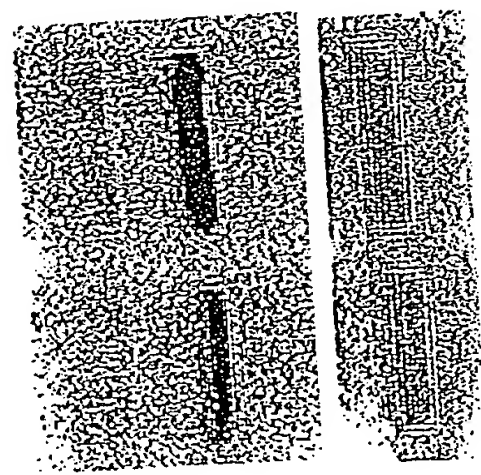
Figure 7F (part 1)

FO6F20"26680660

F

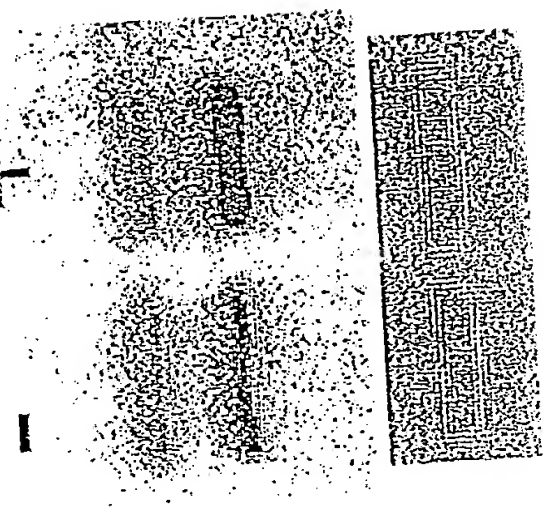
Th2
HAE4A6

+



Th2
CDC 35

+



α CD3

- mTid-1L

- mTid-1S

- actin

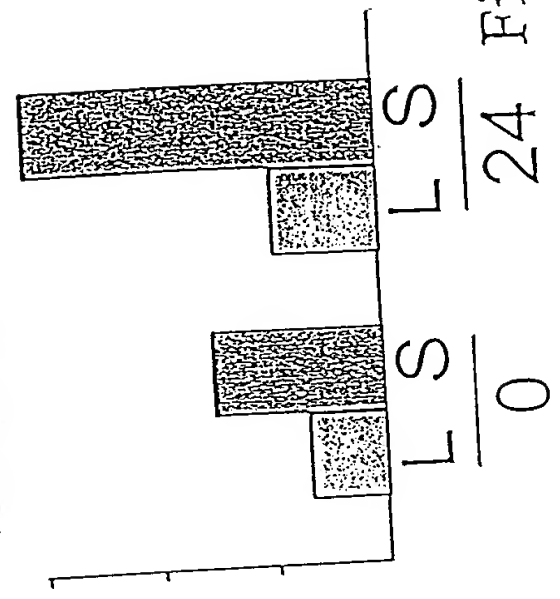
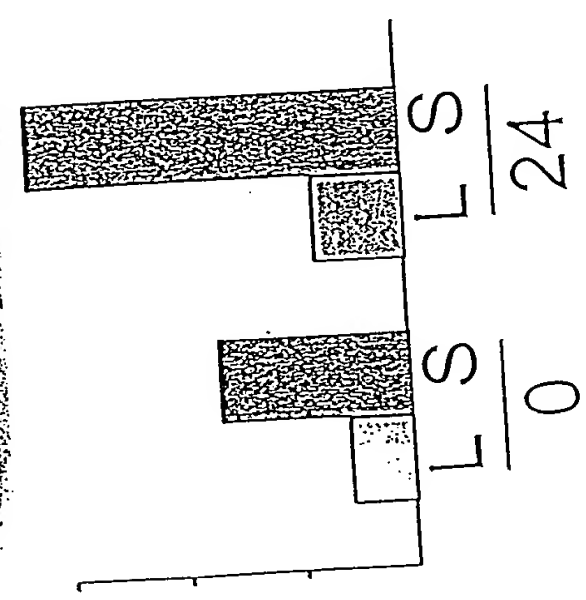


Figure 7F (part 2)

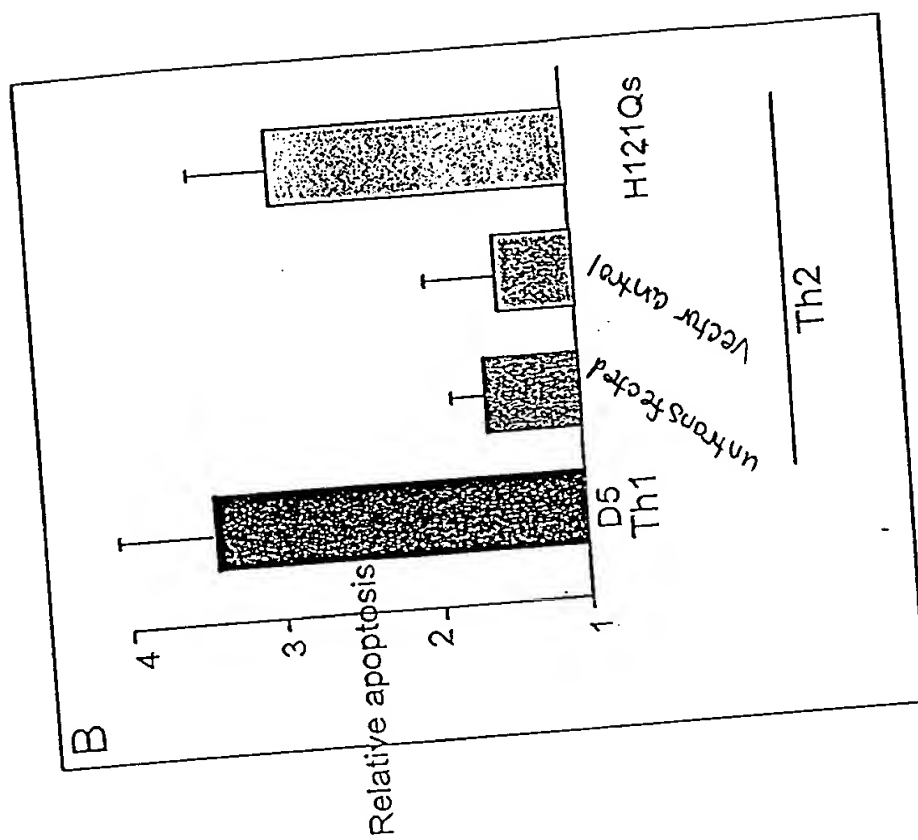


Figure 8

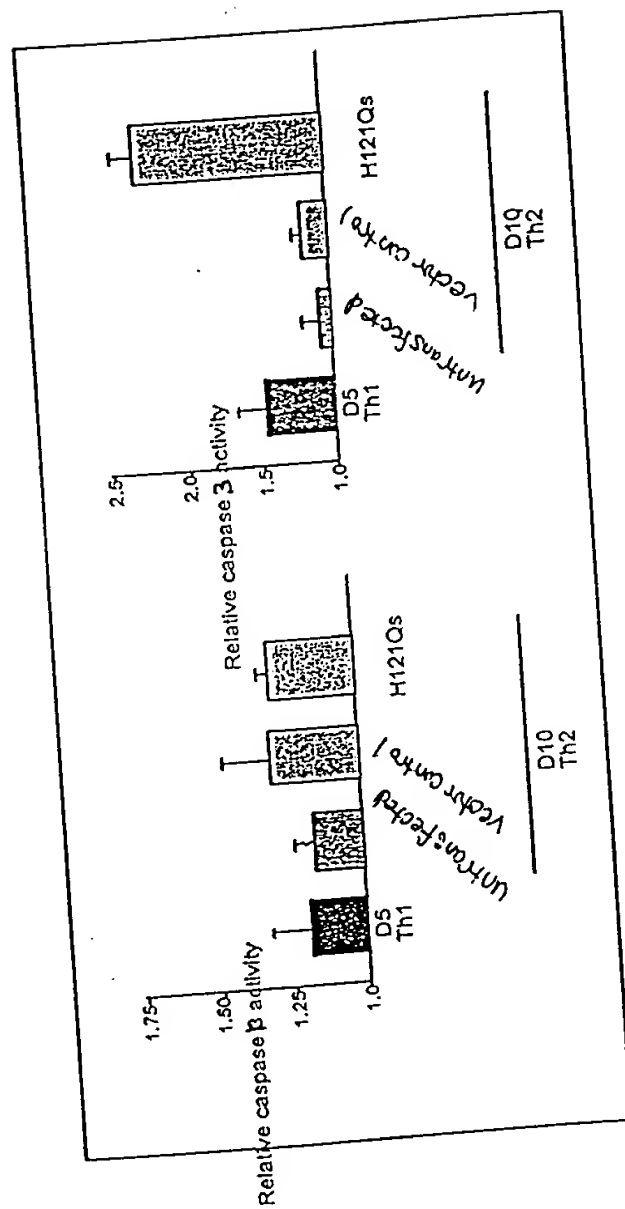


Figure 9

Figure 10

Amino acid sequence of mTid-1L:

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KKAYYQLAKKYHPDTNKDDPKAKEKFSQLAEAYEVLSDEVKRKQYDAYGSAGF
DPGTSSSGQGYWRGGPSVDPEELFRKIFGEFSSSPFGDFQNVVXXXXXXXXXXXXX
XXXXXXXXKEFTVNIMDTCERC DGKGNEPGTKVQHCHYCGGSGMETINTGPFVMR
STCRRCGGRGSIITNPCVVCRGAGQAKQKKRVTIPVPAGVEDGQTVRMPVGKREIF
VTFRVQKSPVFRRTCADIHSDLFISIAQAILGGTAKAQGLYETINVTIPAGIQTDQKI
RLTGKGIPRINSYGYGDHYHIKIRVPKRLSSRQQNLILSYAEDETDVEGTVNGVTH
TSTGGRTMDSSAESKDRREAGEDNEGFLSKLKKIFTS

Amino acid sequence of mTid-1S:

MAAWCSPRWLRVAVGTPRLPAAAGRGVQQPQGGVVATSLCRKLCVSAFGLSMG
AHGPRALLTLRPGVRLTGTKSFPFVCTTSFHTSASLAKDDYYQILGVPRNASQKDI
KKAYYQLAKKYHPDTNKDDPKAKEKFSQLAEAYEVLSDEVKRKQYDAYGSAGF
DPGTSSSGQGYWRGGPSVDPEELFRKIFGEFSSSPFGDFQNVVXXXXXXXXXXXXX
XXXXXXXXKEFTVNIMDTCERC DGKGNEPGTKVQHCHYCGGSGMETINTGPFVMR
STCRRCGGRGSIITNPCVVCRGAGQAKQKKRVTIPVPAGVEDGQTVRMPVGKREIF
VTFRVQKSPVFRRTCADIHSDLFISIAQAILGGTAKAQGLYETINVTIPAGIQTDQKI
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TSTGKRSTGN

106120-2660660

0990892-071901

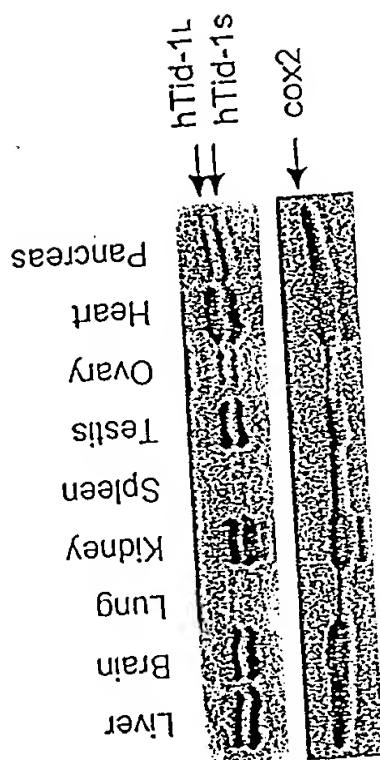


Figure 11

Expression of the TID1 proteins in different tissues -
cox2 is a marker for mitochondria